# Biodosimetry for long-term low-dose past radiation exposure and

What is the lowest detectable dose?

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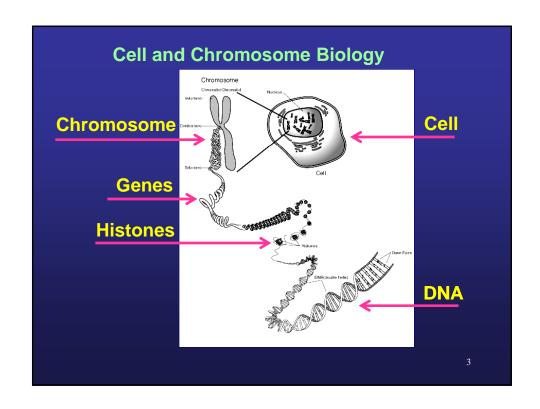
May 18, 2011

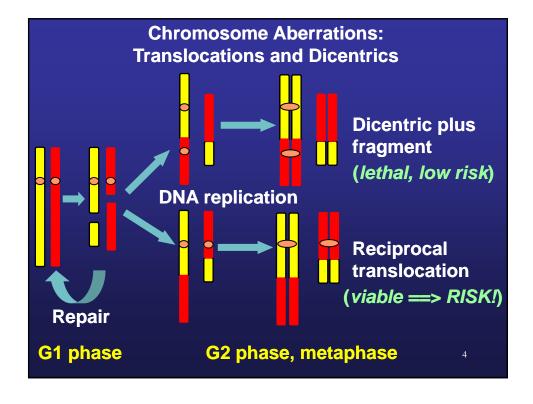


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#### **Outline of this Talk**

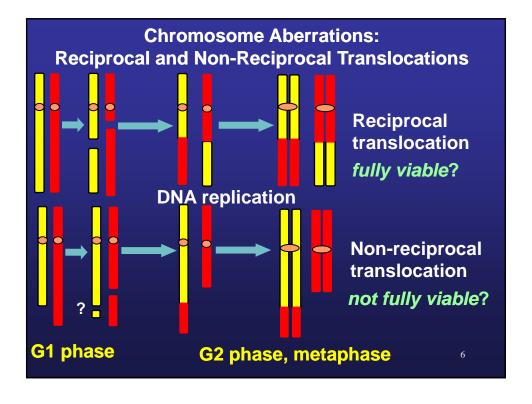
- Chromosome aberrations
  - unbanded
  - banded karyotyping
  - painting
- Dosimetry confounders lessons learned
- Biodosimetry and translocation persistence
- What is the lowest detectable dose?

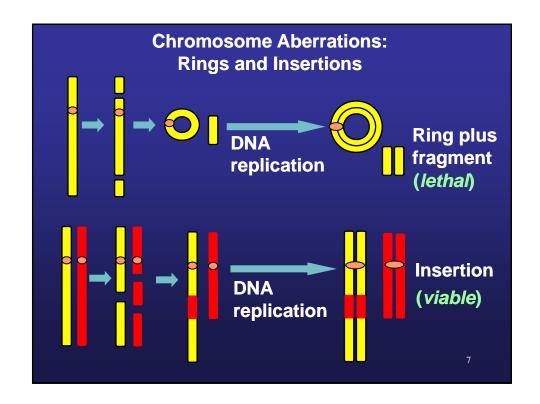


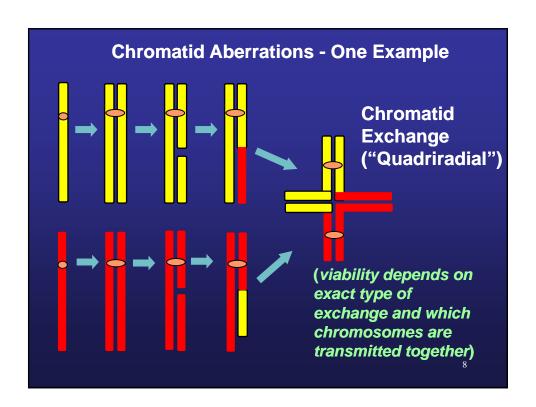


#### **Characteristics of Translocations**

- Induced at frequencies equal to dicentrics
- Stable through cell division
  - persist in vivo indefinitely
  - whereas dicentrics disappear rapidly
- Dosimetry for acute exposure is known
- Accumulate with chronic exposure
- Ideal for biodosimetry ("Gold Standard")
  - chronic exposure
  - exposure occurred many years previously

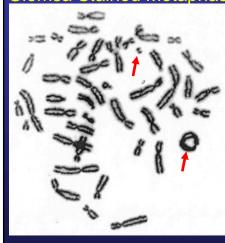






#### **Unbanded Chromosomes**

**Giemsa Stained Metaphase** 



- Detects "unstable" events
- Used widely in research
- Moderate analysis speed
- Inexpensive reagents

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#### **Chromosome Aberrations - Unbanded**



Human cell in metaphase stained with Giemsa

With conventional stains, some categories of chromosome aberrations cannot be seen.

Fragments - yes

Dicentrics - yes

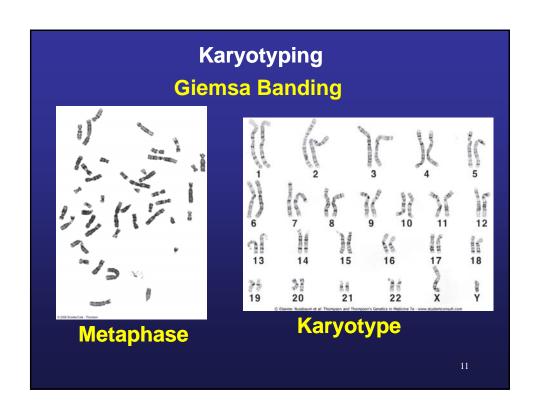
Chromatid damage - yes

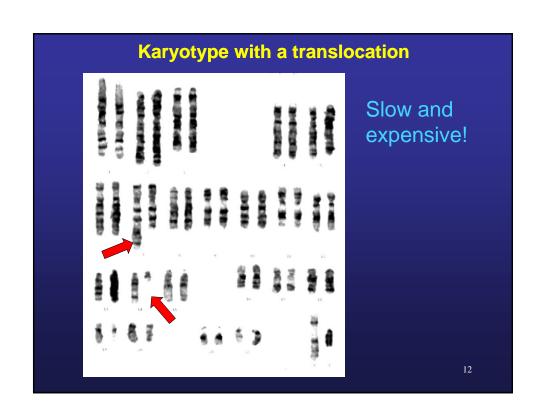
Translocations - generally no

Insertions - no

Complex rearrangements - no

Resolution is limited!





# Karyotyping

With chromosome banding, all categories of chromosome aberrations can be seen.

Fragments - yes

Dicentrics - yes

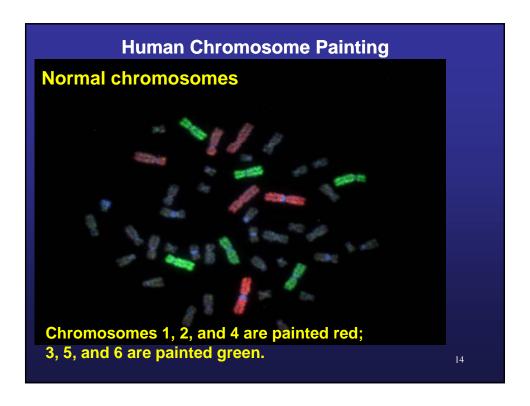
Chromatid damage - yes

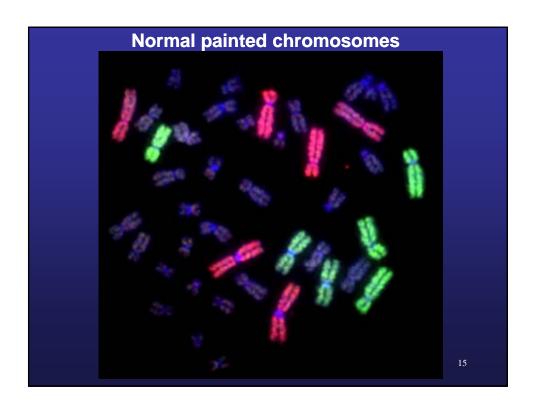
Translocations - yes

Insertions - yes

Complex rearrangements - yes

Speed is limited!



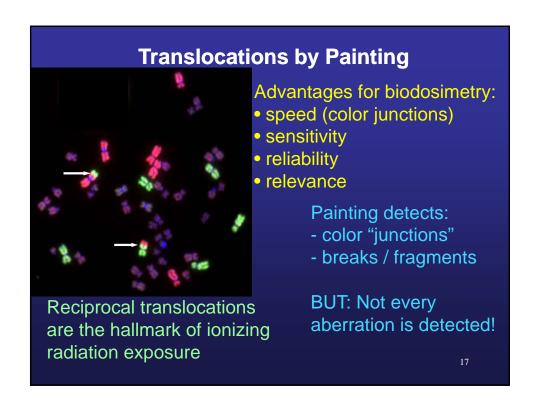


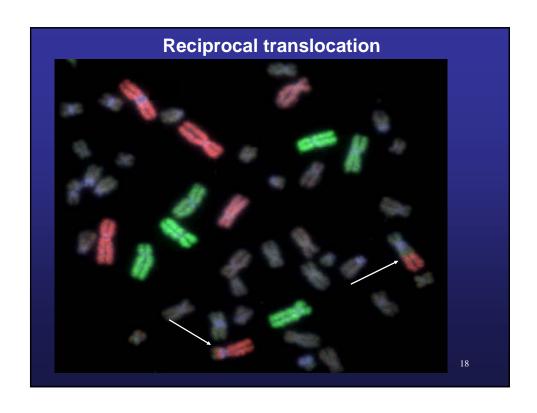
What is the difference between FISH and Chromosome Painting?

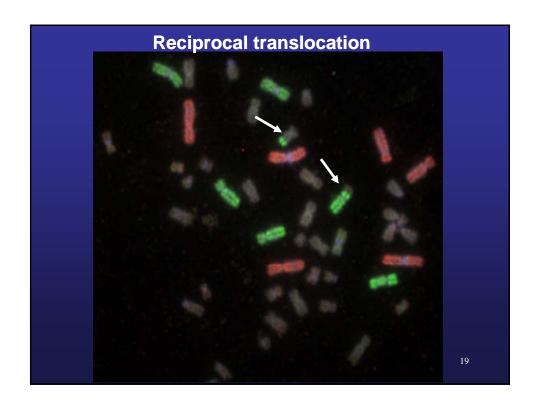
FISH: fluorescence in situ hybridization

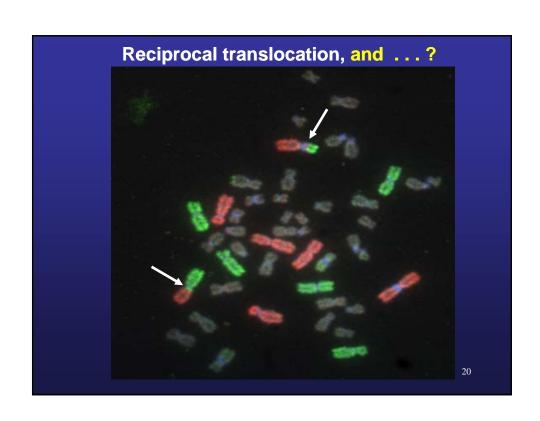
**Chromosome painting: one of many applications of FISH** 

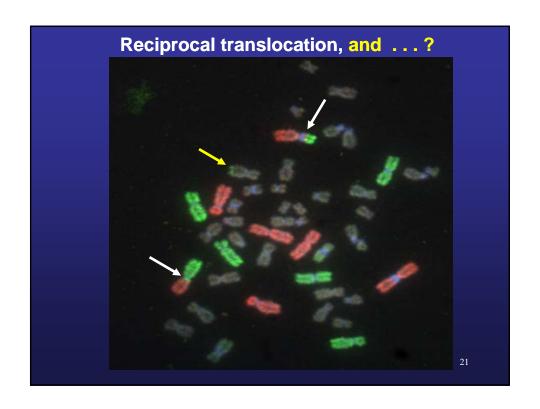
Not all chromosome painting is done by FISH Not all FISH is chromosome painting

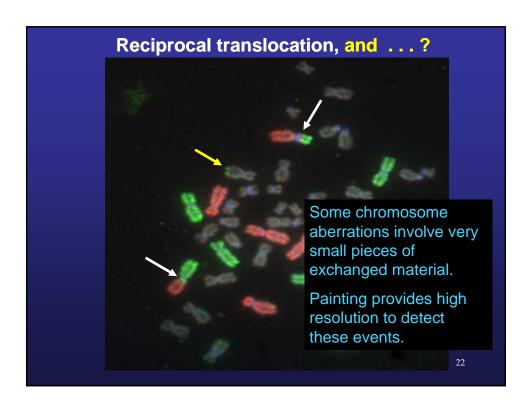


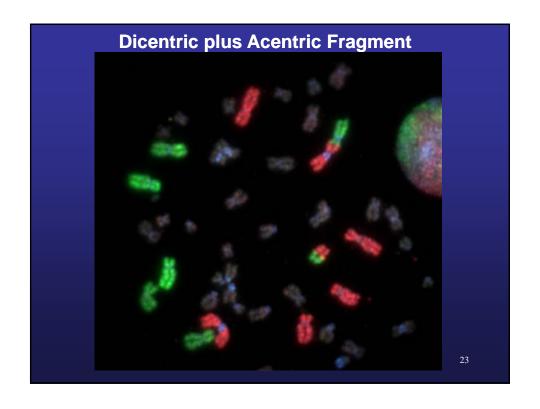


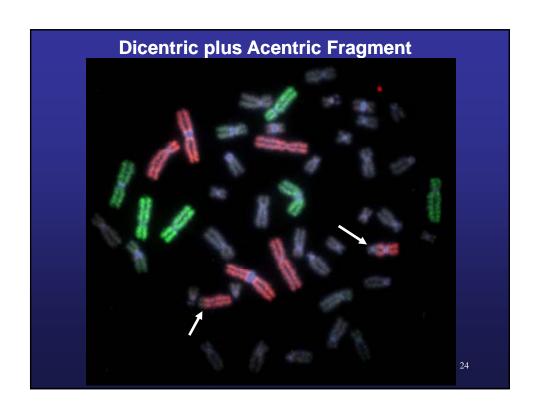


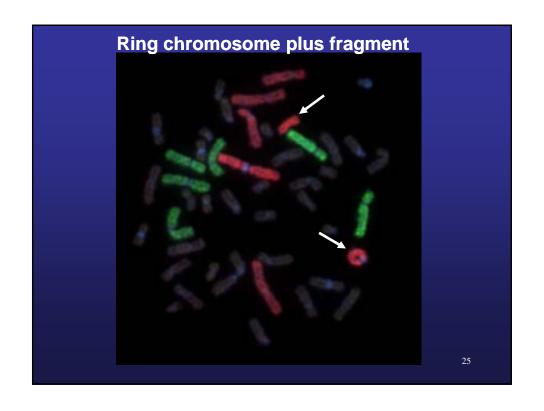


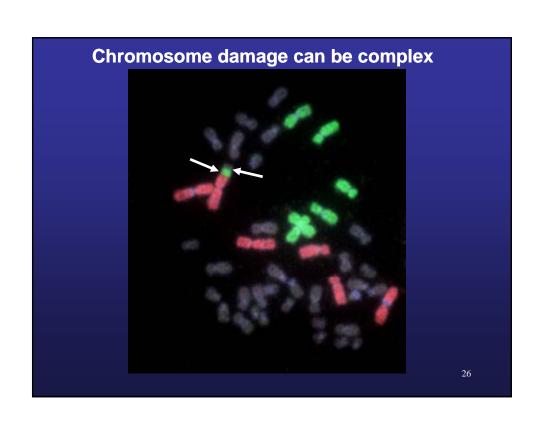


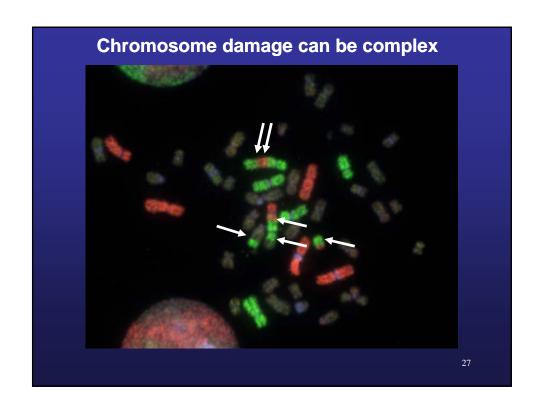




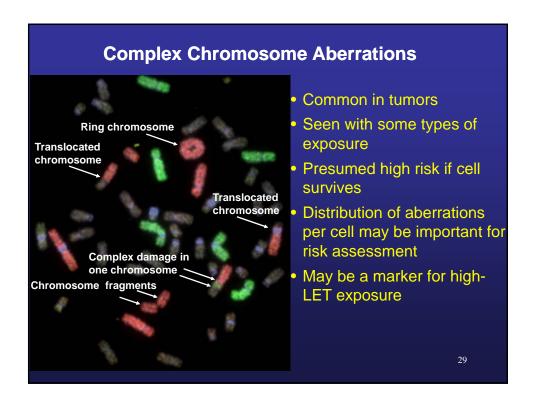


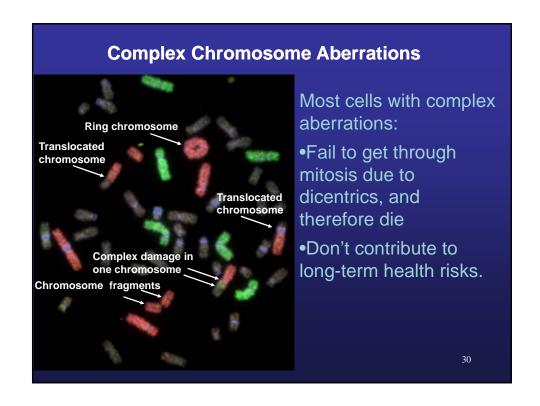












#### Cell Equivalents (CEs), Whole Genome Equivalents

Interchangeable terms.

Chromosome painting does not detect every exchange (i.e., translocation, dicentric) in a given cell.

Detectable exchanges have color junctions.

Undetectable exchanges are those that occur between chromosomes labeled in the same color.

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#### Cell Equivalents (CEs), 1 color painting

Let p = fraction of the genome that is painted Let q = fraction of the genome that is not painted (counterstained, usually in blue)

Then p + q = 1, and  $p^2 + 2pq + q^2$ 

p<sup>2</sup> = unions of broken ends that were both painted

 $q^2$  = unions of broken ends that were both <u>un</u>painted

2pq = unions between one painted and one unpainted chromosome. *This is the observable fraction of all exchanges.* 

#### Cell Equivalents (CEs), 2 color painting

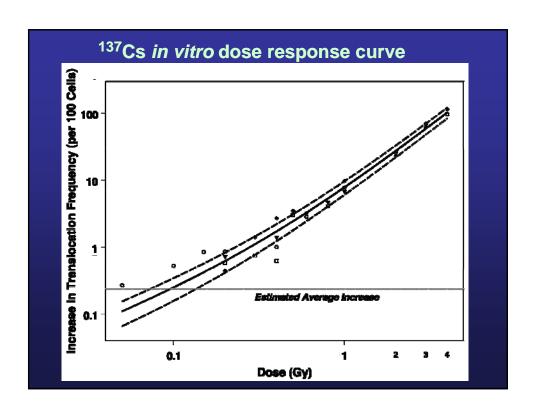
With two color painting,

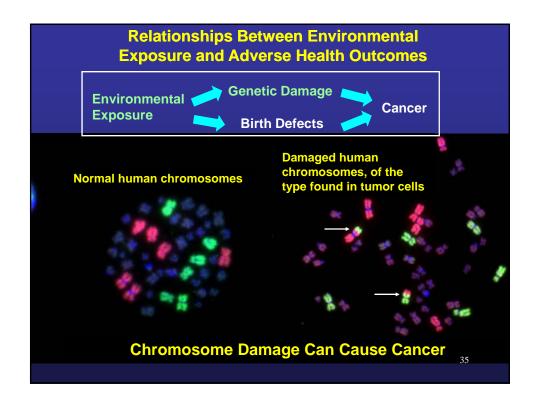
$$p + q + r = 1$$
  
 $p^2 + 2pq + q^2 + 2pr + 2qr + r^2$ 

Detectable color junctions are 2pq + 2pr + 2qr

Need to know which chromosomes are painted, and the percent of the genome they represent.

Reference: Tucker, J.D. (2010) Environmental and Molecular Mutagenesis 51:815:824.







#### Parameters for Radiation Exposure Assessment

<u>Induction</u> - is a measure of the relationship between exposure (dose) and some type of genetic response.

<u>Persistence</u> - is a measure of the longevity of induced damage.

<u>Accumulation</u> - is a measure of the total amount of damage in a cell, tissue, animal or person. Combines induction and persistence.

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#### **Principles for Retrospective Exposure Analysis I.**

- 1. Selection against cells damaged by exposure does not occur or can be taken into account.
- 2. Translocation frequencies pre-existing in the exposed individuals should be known or be estimated from appropriate controls.
- 3. Clones of cytologically abnormal cells are recognizable, and their number and prevalence can be accurately measured.

#### Principles for Retrospective Exposure Analysis II.

- 4. Breaks are distributed among chromosomes in a manner that is proportional to their size.
- 5. The rate of exposure is known, and the effects of dose rate upon translocation frequencies are understood.
- 6. The influence of other confounding exposures, which may fluctuate with time, are negligible.
- 7. The importance of recent exposure history for determining subsequent biological responses, *i.e.*, "adaptation," is known.

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#### **Principles for Retrospective Exposure Analysis III.**

- 8. Tumor cells are not present in the tissue being analyzed.
- 9. Changes in the frequency of genetic damage with age must be well characterized.
- 10. Susceptibility to radiation-induced chromosome aberrations is independent of age.
- 11. Differences between individuals with respect to the above considerations are negligible, or we can adjust for them.

To the extent these principles hold true, dosimetry using translocations can be achieved many years after exposure.

# Dosimetry Confounders Lessons Learned from Human Studies

Very important

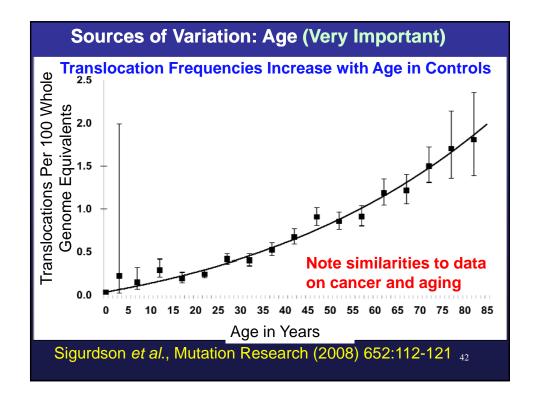
age genotype

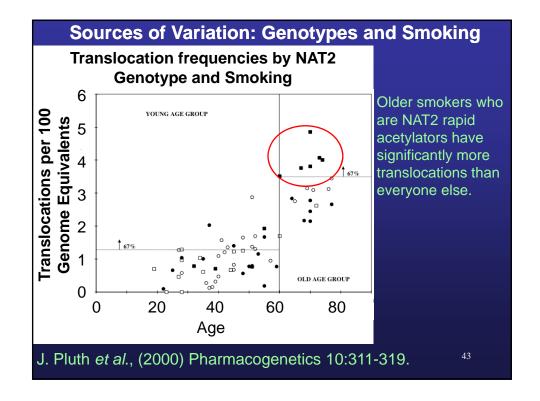
time since exposure

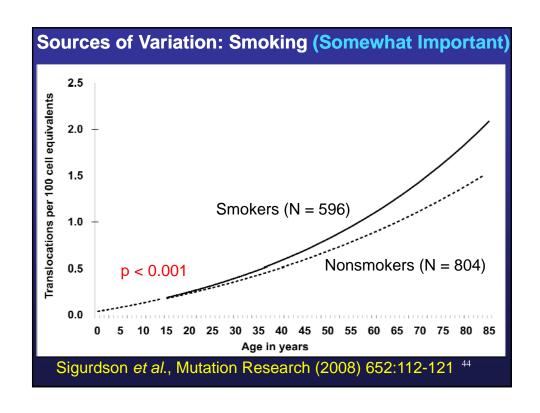
Somewhat important

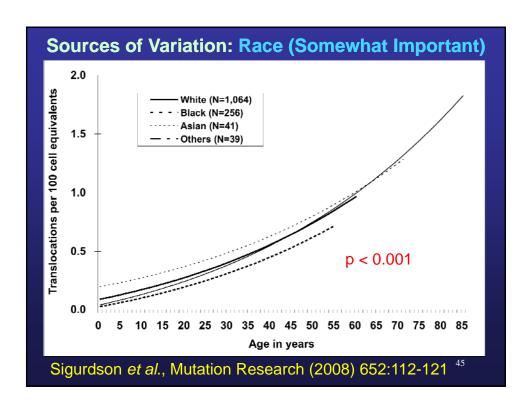
smoking (depends on amount smoked) race (depends on the study)

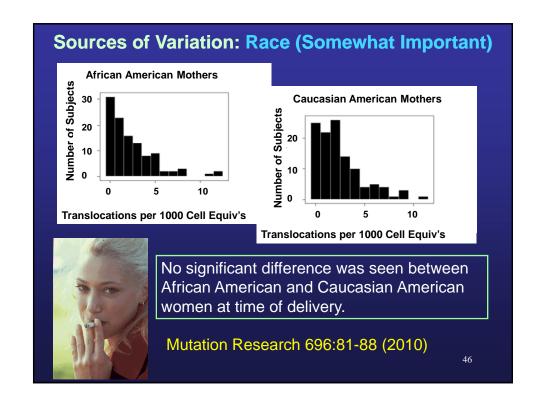
Not important gender

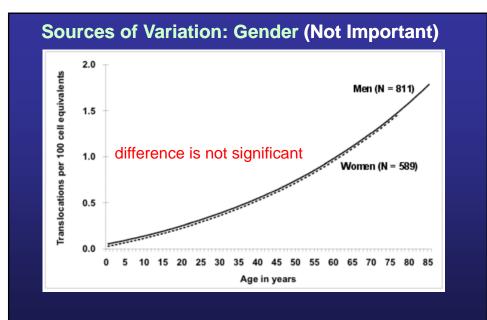












Sigurdson et al., Mutation Research (2008) 652:112-121 47

#### **Sources of Variation**

<u>Genotype</u>: Probably quite important, but the effects of individual genes and alleles are difficult to quantify.

<u>Smoking</u>: Results vary from study to study. Appears to be important for estimating risk, depends on amount smoked (pack years or similar metric).

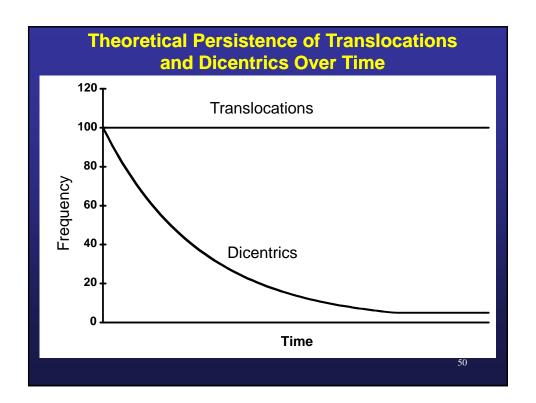
<u>Race</u>: Results vary. Only a few studies done. May be important for estimating risk but more work needs to be done.

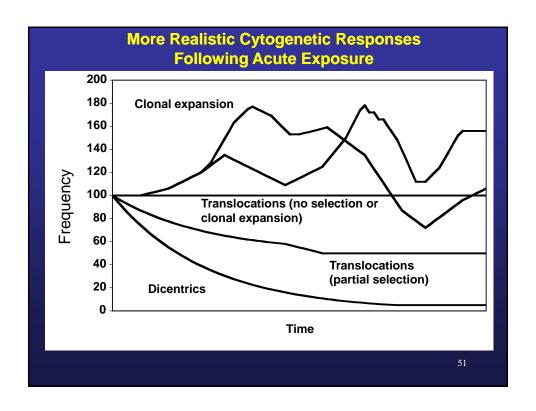
Gender: Not shown to be important.

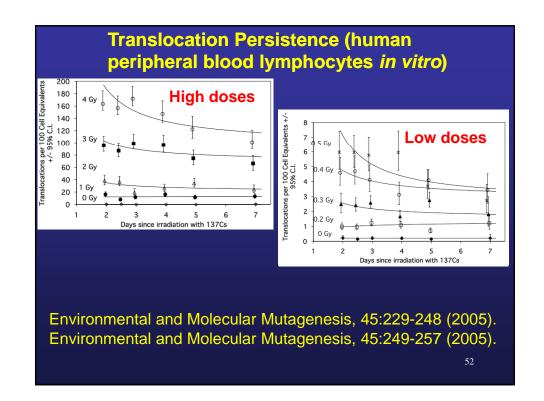
#### **Dosimetry Confounders – Time Since Exposure**

One key assumption for retrospective dosimetry is that translocations *persist*.

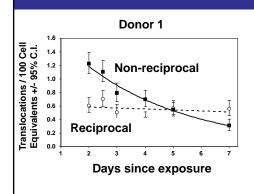
This makes translocations ideal for assessing temporally-displaced exposure.

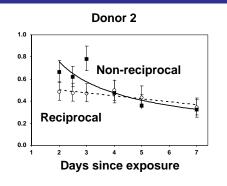












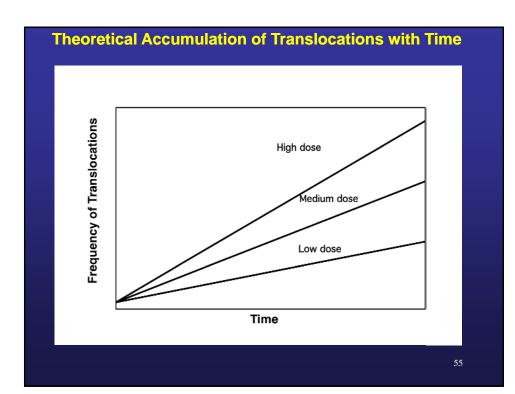
\* human peripheral blood lymphocytes in vitro

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#### **Dosimetry Confounders – Duration of Exposure**

The <u>second</u> key assumption is that translocations *accumulate* with exposure.

Translocations are thus ideal for assessing chronic exposures.

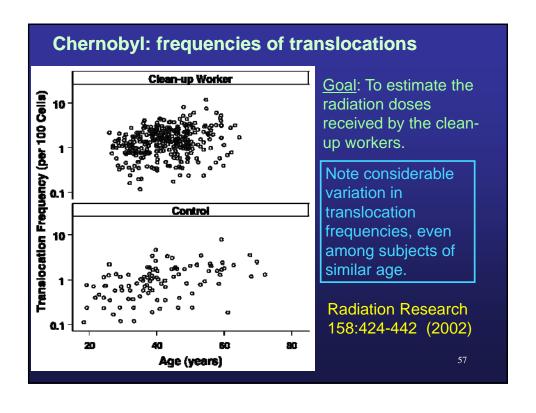


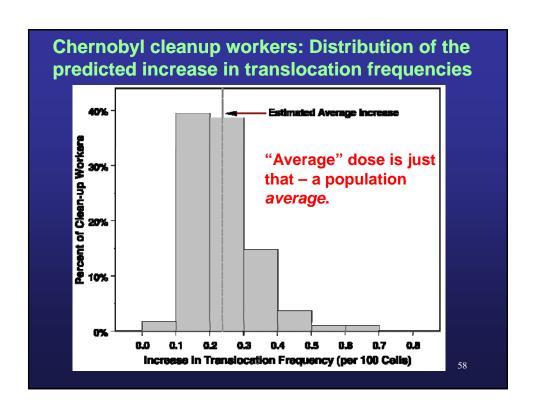
# Why is Understanding Translocation Persistence so Important?

Translocations *are* lost with time but eventually reach a plateau that is greater than the original baseline

Ability to perform dosimetry is retained

Understanding the kinetics of translocation loss is important for accurate dosimetry





#### **Radiation Genotoxicity from Chernobyl**

#### Results:

- 1. The average dose to the clean-up workers was ~9.5 +/- 2.2 cGy, which is half the anticipated dose.
- 2. Translocation frequencies increase significantly with age and smoking.
- 3. Cytogenetic analyses have the power to detect a radiation exposure effect in the presence of confounding factors.

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What about dosimetry for individuals?

How low can FISH biodosimetry go?

#### Low Dose Biodosimetry: Importance of Controls

When estimating exposure, accurate translocation counts from control(s) are essential

- same donor: ideal, usually not possible
- matched control: practical, but limits detection power
- population reference: requires control population

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# Low Dose Biodosimetry: What is the lowest detectable dose?

The answer depends on age and whether the dose is acute or chronic. Other major issues are:

- smoking status
- control sample matching
- time since exposure
- radiation quality
- level of effort expended (counting statistics)

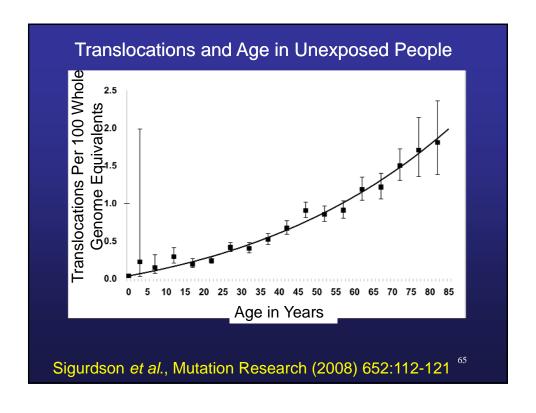
# Two possible biodosimetry scenarios for an exposed individual

- 1. Pre-exposure sample is available
- 2. Pre-exposure sample is not available

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#### **Assumptions**

- Calculations are for a putatively exposed individual, not for a population.
- Pre-exposure sample is not available.
- · Historical data are used for controls.
- Sensitivity to ionizing radiation does not change with age.

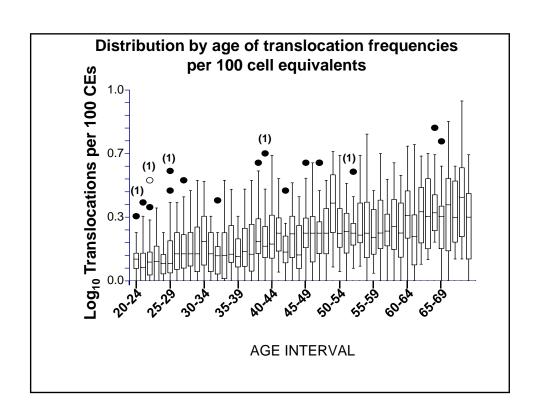


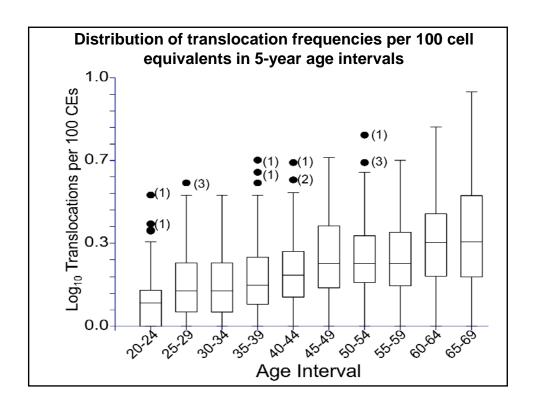
Summary of the Sigurdson <i>et al</i> . Data												
	age interval	N	per 100 CEs	stdev								
	0 - 4	299	0.04	0.09								
N = 1933 subjects.	5 - 9	38	0.16	0.25								
	10 - 14	29	0.22	0.26								
Allana ann anamali.	15 - 19	65	0.21	0.25								
All were apparently	20 - 24	191	0.31	0.34								
normal, healthy	25 - 29	177	0.53	0.54								
people.	30 - 34	138	0.53	0.51								
	35 - 39	154	0.65	0.61								
None had been treated	40 - 44	141	0.74	0.61								
with radiation or	45 - 49	152	1.06	0.83								
	50 - 54	140	1.02	0.80								
chemotherapy.	55 - 59	112	1.00	0.75								
	60 - 64	87	1.41	1.09								
"Control" subjects in	65 - 69	102	1.49	1.27								
other studies.	70 - 74	61	1.66	1.02								
	75 - 79	22	1.86	1.16								
	80 - 85	25	2.17	1.47								

## Step 1 in our analyses

Using the data of Sigurdson *et al.* (2008), we calculated the number of translocations per 100 whole-genome cell equivalents (CEs) in one and five-year age intervals from birth to 80+.

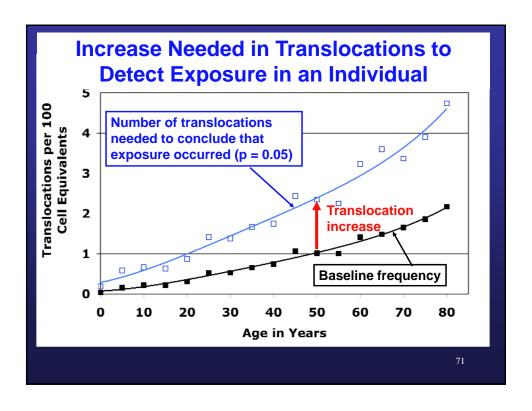
Outliers were removed to satisfy the assumptions of the regression analyses.





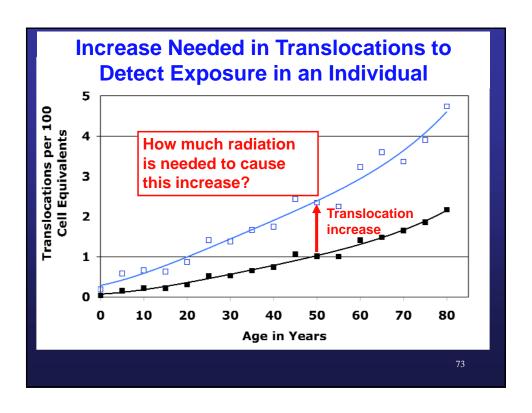
### Step 2 in our analyses

For each 5-year age interval, we calculated the number of translocations per 100 CEs needed in a putatively exposed individual to conclude that a significant increase had occurred with probabilities p = 0.05, p = 0.01, and p = 0.001.



## **Step 3 in our analyses**

For each 5-year age interval we calculated the dose that is needed to induce the minimum number of translocations required for a statistically significant increase.



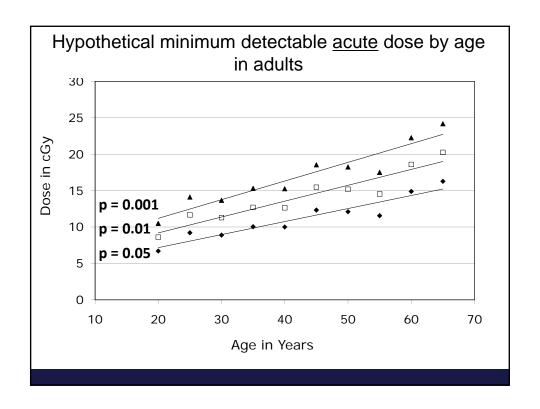
### Step 3, concluded

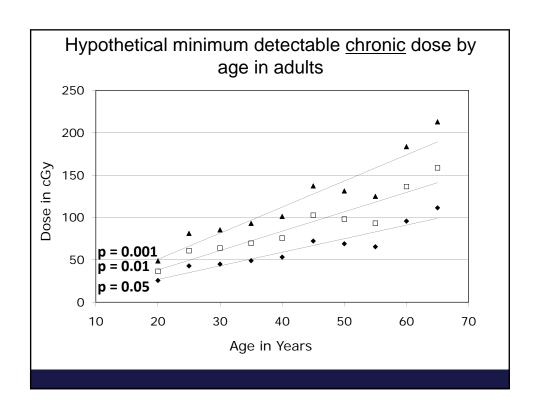
We used the dose response curve generated *in vitro* with <sup>137</sup>Cs (Jones *et al.*, 2002, *Rad. Res.* 158, 424-442).

Number of translocations per CE =  $k + 0.019D + 0.0597D^2$ 

#### where:

k is the baseline translocation frequency D is the dose in Gy



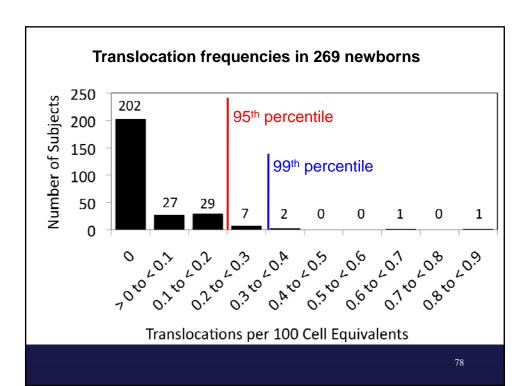


# Parameters of the regression fits

Table 2. Parameters of the regression fits.

		slope			intercept			
Exposure duration	Significance level evaluated	coefficient	standard error	p-value	coefficient	standard error	p-value	Adjusted R- squared
Acute	0.05	0.179	0.022	< 0.0001	3.592	0.969	0.006	0.88
	0.01	0.218	0.026	< 0.0001	4.842	1.176	0.004	0.88
	0.001	0.256	0.031	< 0.0001	6.073	1.373	0.002	0.88
Chronic	0.05	1.591	0.265	< 0.0001	-7.55	11.87	0.54	0.80
	0.01	2.270	0.377	< 0.0001	-11.03	16.92	0.53	0.80
	0.001	3.055	0.508	< 0.0001	-15.38	22.77	0.52	0.80

Tucker, J.D. and Luckinbill, L.S. (2011) Radiation Res. 175:631-7



#### Summary – lowest detectable dose

The dose needed to cause a statistically significant elevation in translocations in an individual increases linearly with age

Acute exposure: 0.18 cGy per year of age

Chronic exposure: 1.59 cGy per year of age

Detecting a given level of exposure is more challenging in older than in younger people.

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### Summary of FISH Radiation Biodosimetry

#### Translocations are the preferred endpoint

- FISH painting is the best method
- fast and reliable
- time since exposure important but not essential
- sensitive enough to detect low doses
  - ✓ populations
  - √ individuals

#### Major confounders:

- age
- cigarette smoking
- possibly genotype (requires more research)

